

Fig.1

10 20 30 40 50 60  
 AACGGATCTG CCCGGCTCAG CCTCCCAAAG TGCTGGGATT GCAGGCGTGA GCCACCTCAC  
 70 80 90 100 110 120  
 CTGGGTACAA GTTTTCAAAA TACATTTATC TAGTACCCAT ACATTCTCCA GTTTGTCCAC  
 130 140 150 160 170 180  
 AGGACATCTT ATGACTTGAG CAAGCTGCTA AAAATCCAAG GGTGCAGCGT TTGTATGTCT  
 190 200 210 220 230 240  
 ATAGGATTGC TCAGACTCTGC CCCCACCCCTG AAAGAATTTA AGAGAATTTT TTGAGGCCAG  
 250 260 270 280 290 300  
 GCACAGTGGC TCACACCTGT AATTCCAGTA CTGTGAGAGT CCGAGGTCAG AGGACTGCTT  
 PPPE  
 310 320 330 340 350 360  
 GAGGCCAGGA GTTCAAGAGC AGCCTGGACA ACACAGGGAG ACCTGTCACT ACAAAGAATA  
 370 380 390 400 410 420  
 AATAAATTAG CCAGGCTTAG TGGCTCATCC CTGTGGTCCC AGCTACTAGG GAGGCAGAAG  
 430 440 450 460 470 480  
 TAGGACTGTG TGTCACAGGA GGTCAAGACT GCAGTGAGCT GAGACCCAGC CACCTGCATT  
 490 500 510 520 530 540  
 CCAGCCTGGG CAACAAAAAG AGACCCTGTC TCAAAAAATA AGTTAAATAA ATAAATAATA  
 550 560 570 580 590 600  
 AAAATAGTTT AAACCTTAAA CACATCTTCT TTTTCAAAGA GGACTTCTTA AGGACTTCAT  
 610 620 630 640 650 660  
 GCTGCGTCTT GTTGATCTCC ACTTCCCTTT TTCAGCGTCC AACTTTTAA CAGTCTCTTT

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670	680	690	700	710	720
TGCCAAGGAT	AATAAGTATA	TAGTTTCTGG	AATCCAGATT	CTTCCCTGTT	TGGACAGCCA
730	740	750	760	770	780
GGGGGACAAT	TTTTGGTCTG	CAGGCCTTTG	<u>CATCTGTTCT</u>	GCTGTTGCTC	AGCAATCTCA
			GRE		
790	800	810	820	830	840
CAGCAAATTT	GCCGAGCCTC	TCCGGAATGC	ACAGCCAGAC	AGAGCTCAGC	GCAAAAGCTA
850	860	870	880	890	900
GAGAACCTGG	CGGAGGGAGA	CTCACAGTGC	CACAAAAAAA	CTTTATCTTT	TCITTTTTTT
910	920	930	940	950	960
TTTCTTTTCT	TTCTTTCTCT	TTCTTTCTTG	TCTTTCTGTC	TTTCTCTCTC	CTCTCTCTGT
970	980	990	1000	1010	1020
CTTTCITTCC	TCCTTTCITT	TCITTTTTCC	TACATGGCAA	GATCTCCCTCA	TGGCAGAAAT
1030	1040	1050	1060	1070	1080
<u>AACTCGCCTT</u>	GACTTCTGTT	TCCACGCTGC	TTCTGCCAGG	ACCATGCGCT	CGGCGTGTTT
GRE					
1090	1100	1110	1120	1130	1140
TTCTTTCCGC	TATAATTATC	CAGGCCCATC	CCAGCTCTGG	TCCCCTCAGC	TGTTCCCTGG
1150	1160	1170	1180	1190	1200
CAGTCCCTTC	TGCTGGTGAA	AACACATATG	GCGCCGGCCT	GACCAGGGTG	TAAGTGTTGT
1210	1220	1230	1240	1250	1260
AAATATCAGGA	AGATGACTGA	ACGTCTTGGG	GACTCCGTTT	CCTCATTTGA	AAATGGAGGT

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1370	1380	1390	1400	1410	1420	1430	1440
TAATACCAGC	CTTCTTCTAC	TCCCCAAACG	CACGTGTTTG	TCCCGGCCAG	AGGGCCCAAT		
					C/EBP		
1330	1340	1350	1360	1370	1380		
TGTTGGCTGT	TCACGCATCA	GTTACCCCCA	CAGGACGGGT	CAGCCAATTA	AAGGCCAACC		
				C/EBP			
1390	1400	1410	1420	1430	1440		
AGGCCCGGTC	CATCTCTGA	CGCCTTTTCT	CATCCCAGGG	CTGGACAGGC	AGCTGGCCTG		
				MyoD			
1450	1460	1470	1480	1490	1500		
GGCCCCGGCTC	TGCTTTGTCA	CGTGCGGGGG	CCGGCCCGTT	TGCTTGCTCTG	TGTGTAGGAG		
	GRE						
1510	1520	1530	1540	1550	1560		
CGTGAGGTCA	CGCTGGGTGC	TCCCGCCCCG	CCGGGGCCCT	TAGTGTCCCT	GGTCCTTAAA		
1570	1580	1590	1600	1610	1620		
CGCCAGGCCG	CTCCACCGGG	GGAGAAGGCG	CGAAGCCCGAG	CCGAGGCCAA	CGGCTGTTGT		
1630	1640	1650	1660	1670	1680		
CGGTTCGGCG	GCCACCTGTT	GCTGCAGTTC	TGATTGGTTC	CTTCCCCCGA	CAACGCGCGG		
1690	1700	1710	1720	1730	1740		
GCTGTAACCA	ATCGACACGC	AGGCCGGTGC	CGAGGCCCCA	GTCCCGCCCT	GCAGGAGCCA		
	C/EBP						
1750	1760	1770	1780	1790	1800		
GCCCGCGGCT	CGCTCGCAGG	AGGCTGGGTA	GTTTGCCCGAG	CGTAGGGGGG	CTGGGCCCAT		
1810	1820	1830	1840	1850	1860		
AAAAGAGGAA	GTGCACTTAA	GACACGGCCC	CGCTGGACGC	TTGTTAGAAA	CCGTCTGGC		
1870	1880	1890	1900	1910	1920		
TGGGAAGGCA	AGAGGTGTGT	GACTGGACAA	GACTTGTTTC	TGGCGGTACG	TCTTGCCATC		